OIPE

RAW SEQUENCE LISTING DATE: 01/15/2002 PATENT APPLICATION: US/10/027,450 TIME: 15:31:38

Input Set : A:\BB1126 USDIV Seq List.txt
Output Set: N:\CRF3\01152002\J027450.raw

Does Not Comply 3 <110> APPLICANT: Falco, Saverio Carl Corrected Diskette Needed Hitz, William D. 5 Kinney, Anthony J. Cahoon, Rebecca E. Rafalski, J. Antoni 9 <120> TITLE OF INVENTION: PLANT BRANCHED CHAIN AMINO ACID BIOSYNTHETIC ENZYMES 11 <130> FILE REFERENCE: BB-1126 C--> 13 <140> CURRENT APPLICATION NUMBER: US/10/027,450 C--> 14 <141> CURRENT FILING DATE: 2001-12-20 16 <150> PRIOR APPLICATION NUMBER: 60/063,423 W--> 17 <151> PRIOR FILING DATE: 1997 October 28 1997-/0-28 19 <160> NUMBER OF SEQ ID NOS: 54 21 <170> SOFTWARE: Microsoft Word Version 7.0A

ERRORED SEQUENCES

2545 <210> SEQ ID NO: 54 2546 <211> LENGTH: 113 2547 <212> TYPE: PRT 2548 <213> ORGANISM: Lactococcus lactis 2550 <400> SEQUENCE: 54 2551 Met Lys Thr Tyr Lys Gly Thr Ser Val Val Met Asn Asp Asn Asp Thr 2554 Asp Lys Lys Ala Asp Lys Lys Gly Gly Lys Asn Tyr Trp Arg Tyr Lys 20 25 2557 Asp Tyr Asp Asn Asp Asn Ala Lys Tyr Lys Lys Ala Ser Ser Gly Asp 2558 35 40 2560 Asn Gly Ser Gly Ser Ser Arg His Ala Ala Trp Ala Ser Asp Tyr Gly 50 55 2563 Arg Ala Ala Gly Ser Tyr Ser Asp Tyr Asn Asn Ala Lys Asn Gly Lys 70 2564 65 75 2566 Arg Val Asn Thr Lys Ser Ser Thr Asp His Thr Ser Gly Asp His Asp 85 90 2569 Trp Lys Asp Lys Asn Gly Asp Asp Gly Thr Tyr Ala Ser Ala Tyr Lys 2570 100 105 2572 Asn E--> 2576 741 E--> 2578/1.Docket Number delite

VERIFICATION SUMMARY
PATENT APPLICATION: US/10/027,450
DATE: 01/15/2002
TIME: 15:31:39

Input Set : A:\BB1126 USDIV Seq List.txt
Output Set: N:\CRF3\01152002\J027450.raw

L:13 M:270 C: Current Application Number differs, Replaced Application Number L:14 M:271 C: Current Filing Date differs, Replaced Current Filing Date L:17 M:256 W: Invalid Numeric Header Field, Wrong Prior FILING DATE:YYYY-MM-DD L:362 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:5 L:362 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 L:402 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:6 L:402 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6 L:694 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:12 L:694 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12 L:695 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:12 L:695 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12 L:696 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:12 L:696 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12 L:697 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:12 L:697 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12 L:731 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:13 L:731 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13 L:1844 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:37 L:1844 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:37 L:1976 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:39 L:1976 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:39 L:1977 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:39 L:1977 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:39 L:1978 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:39 L:1978 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:39L:1979 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:39 L:1979 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:39 $L:2009\ M:258\ W:$ Mandatory Feature missing, <223> not found for SEQ ID#:40 $L\!:\!2009\ M\!:\!341\ W\!:$ (46) "n" or "Xaa" used, for SEQ ID#:40 L:2061 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:42 L:2061 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:42 L:2416 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:51 L:2416 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:51 L:2576 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:54 L:2578 M:333 E: Wrong sequence grouping, Amino acids not in groups! L:2578 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:2 L:2578 M:252 E: No. of Seq. differs, <211>LENGTH:Input:113 Found:115 SEQ:54